

GenCore version 5.1.4.D5_4578
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OM protein - protein search, using SW model

Run on: March 24, 2003, 15:45:24 ; Search time 11.6788 Seconds
(without alignments)
628.181 Million cell updates/sec

Title: US-09-988-971-2_COPY_35_90

Perfect score: 288
Sequence: 1 ATAAVALGSPAGCPAELSLR.....VLSEVSGREYVPSVHAKV 56

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 908470 seqs, 133250620 residues

Total number of hits satisfying chosen parameters: 908470

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database :

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22: /SIDSI/gcgdata/geneseq/geneseq-emb1/AA2001.DAT.*
23: /SIDSI/gcgdata/geneseq/geneseq-emb1/AA2002.DAT.*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length DB	ID	Description
1	288	100.0	210	23	AAO15458
2	288	100.0	261	23	AAO15457
3	284	98.6	248	21	AA842993
4	284	98.6	261	23	AAU91308
5	242	84.0	259	23	AAO15456
6	148	51.4	70	22	ABG05994
7	145	50.3	395	22	AAU31598
8	99	34.4	315	22	AAU31072
9	96	33.3	509	21	AAU94420
10	91.5	31.8	517	22	AB857557

11	90	31.2	61	23	ABB84051
12	90	31.2	471	23	ABB84052
13	90	31.2	505	22	ABP98322
14	90	31.2	513	23	ABB84055
15	89.5	31.1	2415	22	ABB85710
16	88	30.6	126	20	AAW73554
17	88	30.6	346	21	AAV76750
18	88	30.6	346	22	AAE06208
19	88	30.6	508	21	AAE37700
20	87.5	30.4	60	17	AAW07876
21	87.5	30.4	251	21	AAV44450
22	87.5	30.4	533	14	AAK37705
23	87.5	30.4	533	21	AAV44447
24	87.5	30.4	533	21	AAV44449
25	87.5	30.4	533	21	AAV44451
26	87.5	30.4	533	22	AAE84661
27	85.5	29.7	211	22	AAE36685
28	85.5	29.7	214	22	AAE36681
29	85.5	29.7	536	14	AAE39706
30	85.5	29.7	536	23	AAU78678
31	85	29.5	116	22	AAU31071
32	84.5	29.3	543	20	AAV24421
33	84.5	29.3	543	22	ABG10302
34	84.5	29.3	543	22	AAE84663
35	82.5	28.6	541	23	AAU74614
36	82.5	28.6	542	23	ABE97339
37	82.5	28.6	565	22	ABG23778
38	82	28.5	59	20	AAV28669
39	82	28.5	59	22	AAU08731
40	82	28.5	496	22	AAV29668
41	82	28.5	496	22	AAU08730
42	82	28.5	496	22	AAU08734
43	82	28.5	496	22	AAU08735
44	80.5	28.0	1200	21	AAE19313
45	80	27.8	1683	21	AAV71160

ALIGNMENTS

RESULT 1	AAO15458	AAO15458 standard; Protein; 210 AA.
XX	AAO15458;	
XX		
AC	AAO15458;	
XX		
DT	03-OCT-2002 (first entry)	
XX		
DE	Mouse modulator of antigen receptor signalling short isoform protein.	
XX		
KW	Mouse; gene therapy; modulator of antigen receptor signalling; MARS;	
KW	tumour suppressor gene; Src-like adaptor protein; SLAP;	
KW	myeloid malignancy; acute myelogenous leukemia; autoimmune disorder;	
XX	immunosuppression; myeloproliferative disorder; breast cancer.	
OS	Mus sp.	
XX		
PN	W0200242452-A2.	
XX		
PD	30-MAY-2002.	
XX		
PF	26-NOV-2001; 2001WO-CA01662.	
XX		
PR	27-NOV-2000; 2000CA-2324663.	
XX		
PA	(HOSP-) HOSPITAL FOR SICK CHILDREN.	
XX		
PI	Mcglade JC, Loreto MP;	
XX		
DR	WPI: 2002-566564/60.	
DR	N-PSDB; AAL4090.	
XX		
PT	New isolated modulator of antigen receptor signaling protein or its	

PT fragment, useful for treating malignant disorders such as myeloid
PT malignancies, autoimmune disorders and myeloproliferative disorders -
XX
XX
PS Claim 8; Page 78; 110pp; English.
XX
CC The invention comprises the amino acid and coding sequences of modulator
CC of antigen receptor signalling (MARS) proteins. The MARS protein is a
CC putative tumour suppressor gene and exhibits structural and sequence
CC similarity to the Scr-like adaptor protein (SLAP). The MARS DNA and
CC protein sequences of the invention are useful for the treatment of
CC myeloid malignancies (e.g. acute myelogenous leukaemia) autoimmune
CC disorders, immunosuppression, myeloproliferative disorders and
CC malignancies related to the de-regulation of tyrosine kinases (e.g.
CC breast cancer). The present amino acid sequence represents a mouse MARS
CC protein.
XX
SQ Sequence 210 AA;
XX
Query Match 100.0%; Score 288; DB 23; Length 210;
Best Local Similarity 100.0%; Pred. No. 1.6e-29;
Matches 56; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
Qy 1 ATVALGSPFAGPAGPAGLRLGEPPLTVSESDGMWTVLSEVSGREYNIPEVHAKV 56
Db 35 ATVALGSPFAGPAGPAGLRLGEPPLTVSESDGMWTVLSEVSGREYNIPEVHAKV 90
RESULT 2
ID AA015457
XX AA015457 standard; Protein; 261 AA.
XX
AC AA015457;
XX
DT 03-OCT-2002 (first entry)
XX
DE Human modulator of antigen receptor signalling (MARS) protein.
XX
XX Human; gene therapy; modulator of antigen receptor signalling; MARS;
KW tumour suppressor gene; Scr-like adaptor protein; SLAP;
KW myeloid malignancy; acute myelogenous leukaemia; autoimmune disorder;
KW immunosuppression; myeloproliferative disorder; breast cancer.
XX
OS Homo sapiens.
XX
PN WO200242452-A2.
XX
PD 30-MAY-2002.
XX
PF 26-NOV-2001; 2001WO-C0A01662.
XX
PR 27-NOV-2000; 2000CA-2324663.
XX
PA (HOSP-) HOSPITAL FOR SICK CHILDREN.
XX
PI Meglade JC, Loreto MP;
XX
DR WPI; 2002-566564/60.
XX
DR N-PSDB; AAL44089.
XX
PT New isolated modulator of antigen receptor signalling protein or its
PT fragment, useful for treating malignant disorders such as myeloid
PT malignancies, autoimmune disorders and myeloproliferative disorders -
XX
XX Claim 7; Fig 9A; 110pp; English.
XX
XX The invention comprises the amino acid and coding sequences of modulator
CC of antigen receptor signalling (MARS) proteins. The MARS protein is a
CC putative tumour suppressor gene and exhibits structural and sequence
CC similarity to the Scr-like adaptor protein (SLAP). The MARS DNA and
CC protein sequences of the invention are useful for the treatment of
CC myeloid malignancies (e.g. acute myelogenous leukaemia) autoimmune
CC disorders, immunosuppression, myeloproliferative disorders and
CC malignancies related to the de-regulation of tyrosine kinases (e.g.

CC breast cancer). The present amino acid sequence represents a human MARS
CC protein.
XX
XX
SQ Sequence 261 AA;
XX
Query Match 100.0%; Score 288; DB 23; Length 261;
Best Local Similarity 100.0%; Pred. No. 2.1e-29;
Matches 56; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
Qy 1 ATVALGSPFAGPAGPAGLRLGEPPLTVSESDGMWTVLSEVSGREYNIPEVHAKV 56
Db 35 ATVALGSPFAGPAGPAGLRLGEPPLTVSESDGMWTVLSEVSGREYNIPEVHAKV 90
RESULT 3
ID AAB42993
XX AAB42993 standard; Protein; 248 AA.
XX
AC AAB42993;
XX
DT 08-FEB-2001 (first entry)
XX
DE Human OREF2757 polypeptide sequence SEQ ID NO:5514.
XX
XX Human; open reading frame; OREF; detection; cytosolic; hepatocytic;
KW vulnery; antiproliferative; antiparkinsonian; neuroprotective;
KW anticonvulsant; osteoporotic; antiarthritic; immunosuppressant; cardiant;
KW immunostimulant; thrombolytic; coagulant; vasotropic; antidiabetic;
KW hypotensive; dermatological; immunosuppressive; antineoplastic;
KW antiviral; antibacterial; antifungal; antineumatic; antihypertensive;
KW antianemic; gene therapy; cancer; proliferative disorder; hypertension;
KW neurodegenerative disorder; osteoarthritis; graft vs host disease;
KW cardiovascular disease; diabetes mellitus; hypothyroidism; SCID; AIDS;
KW cholesterol ester storage; systemic lupus erythematosus; infection;
KW severe combined immunodeficiency; malaria; autoimmune disorder; asthma;
KW allergy; aplastic anaemia; nocturnal haemoglobinuria; burn; wound;
KW bone damage; cartilage damage; antineoplastic disease; coagulation;
KW thrombosis; contraceptive.
XX
OS Homo sapiens.
XX
PN WO200058473-A2.
XX
PD 05-OCT-2000.
XX
PF 31-MAR-2000; 2000WO-US08621.
XX
PR 31-MAR-1999; 99US-0127607.
XX
PR 02-APR-1999; 99US-0127636.
XX
PR 05-APR-1999; 99US-0127728.
XX
PR 30-MAR-2000; 2000US-0540763.
XX
PA (CUBA-) CUBAGEN CORP.
XX
PI Shinkens RA, Leach M;
XX
DR WPI; 2000-602362/57.
XX
DR N-PSDB; AAC77202.
XX
PT Novel nucleic acids and peptides derived from open reading frame X,
PT useful for treating e.g. cancers, proliferative disorders,
PT neurodegenerative disorders and cardiovascular disease -
XX
XX Claim 11; Page 4693-4694; 5507pp; English.
XX
XX AACT4446 to AACT7606 encode the proteins given in AAB40237 to AAB43397,
CC which represent the human OREF open reading frames 1 to 3161. The OREF
CC sequences have activities such as: cytosolic; hepatocytic; vulnery;
CC antiproliferative; antiparkinsonian; neuroprotective; cardiant;
CC osteoporotic; antidiabetic; antineumatic; antihypertensive;
CC immunostimulant; cardiant; thrombolytic; coagulant; vasotropic;
CC antidiabetic; hypotensive; dermatological; immunosuppressive;
CC antineoplastic; antineumatic; antihypertensive; antineumatic;
CC antineoplastic; antineumatic; antihypertensive; antineumatic;
CC antineoplastic; antineumatic; antihypertensive; antineumatic;

CC antihypertoid, and antianemic. The sequences can be used for determining
CC the presence of or predisposition to, or preventing or treating
CC pathological conditions associated with an ORFX-associated disorder. The
CC nucleic acids can be used to express ORFX proteins in gene therapy
CC vectors. The proteins and nucleic acids may be used to treat cancers,
CC proliferative disorders, neurodegenerative disorders, osteoarthritis,
CC graft vs host disease, cardiovascular disease, diabetes mellitus
CC hypertension, hypothyroidism, cholesterol ester storage, systemic lupus
CC erythematosus, severe combined immunodeficiency (SCID), AIDS, viral,
CC bacterial or fungal infection, malaria, autoimmune disorders, asthma,
CC allergies, aplastic anaemia, burns, wounds, bone and cartilage damage,
CC nocturnal haemoglobinuria, antiinflammatory disease, to enhance
CC coagulation, to inhibit thrombosis, and as a contraceptive.

XX Sequence 248 AA;

Query Match 98.6%; Score 284; DB 21; Length 248;
Best Local Similarity 98.2%; Pred. No. 6,6e-29;
Matches 55; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 1 ATAAAGSPFAGGPAELSLRGEPLTIVSEDDGDMWTIVSEVSGREYNIPSVYAKV 56
DB 22 ATAAAGSPFAGGPAELSLRGEPLTIVSEDDGDMWTIVSEVSGREYNIPSVYAKV 77

RESULT 4
AAU91308 standard; Protein; 261 AA.

XX AAU91308;

DT 18-JUN-2002 (first entry)

XX Human protein NOV13.

XX Human; NOVX; gene therapy; cardiomyopathy; atherosclerosis;
XX cell signal processing disorder; metabolic pathway modulation disorder;
XX diabetes; cancer; adenocarcinoma; lymphoma; prostate cancer;
XX uterine cancer; immune response; graft-versus-host disease;
XX acquired immunodeficiency syndrome; AIDS; asthma; Crohn's disease;
XX hypertension; congenital heart defects; multiple sclerosis; inflammation;
XX Albright hereditary osteodystrophy.

XX Homo sapiens.

XX MO200216599-A2.

XX 28-FEB-2002.

XX 27-AUG-2001; 2001MO-US26510.

XX 25-AUG-2000; 2000US-228191P.

XX 08-FEB-2001; 2001US-267300P.

XX 20-FEB-2001; 2001US-26961P.

XX 20-MAR-2001; 2001US-277337P.

XX (CURA-) CURAGEN CORP.

XX (CORT-) CORT THERAPEUTICS INC.

XX Burgess CE, Conley PB, Grose WM, Hart M, Kekuda R, Shimkets RA;

XX Spletke KA, Szekeres BS, Tomlinson JE, Topper JN, Yang R;

XX WPI; 2002-280937/32.

XX N-PSDB; ABK61465.

XX New polypeptides for treating or preventing a disorder associated with
XX them, in humans, e.g. cardiomyopathy, atherosclerosis or cancers -
XX Claim 3; Page 98; 263pp; English.

CC NOVX is NOV1-14, 15a, 15b, 16a, and 16b. The NOVX polypeptide, nucleic
CC acid encoding it and antibody against it, are useful for treating or
CC preventing (e.g. by gene therapy) a NOVX-associated disorder in humans,
CC e.g. cardiomyopathy, atherosclerosis, a disorder related to cell signal
CC processing and metabolic pathway modulation, diabetes or cancers. The
CC NOVX polypeptide and nucleic acids are also useful for determining the
CC presence of predisposition to the diseases. The NOVX nucleic acid and
CC polypeptide are especially useful in therapeutic or prophylactic
CC applications for disorders associated with aberrant NOVX expression or
CC activity, e.g. cancers (e.g. adenocarcinoma, lymphoma, prostate cancer or
CC uterine cancer), immune response, graft-versus-host disease, hypertension,
CC immunodeficiency syndrome (AIDS), asthma, Crohn's disease, hyperextension,
CC congenital heart defects, multiple sclerosis, inflammation of Albright
CC hereditary osteodystrophy and many other diseases listed in the
CC specification. The DNA encoding the protein is useful in gene therapy
CC for treating the conditions. This is also useful in detection assays, or
CC for chromosome mapping, tissue typing, diagnostic or prognostic assays, or
CC for developing a powerful assay system for functional analysis of
CC various human disorders, as well as in diagnostic applications. The
CC present sequence represents a NOVX protein.

XX Sequence 261 AA;

Query Match 98.6%; Score 284; DB 23; Length 261;
Best Local Similarity 98.2%; Pred. No. 7.1e-29;
Matches 55; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 1 ATAAAGSPFAGGPAELSLRGEPLTIVSEDDGDMWTIVSEVSGREYNIPSVYAKV 56
DB 35 ATAAAGSPFAGGPAELSLRGEPLTIVSEDDGDMWTIVSEVSGREYNIPSVYAKV 90

RESULT 5
AA015456 standard; Protein; 259 AA.

XX AA015456;

DT 03-OCT-2002 (first entry)

XX Mouse modulator of antigen receptor signalling (MARS) protein.

XX Mouse; gene therapy; modulator of antigen receptor signalling; MARS;
XX tumor suppressor gene; Src-like adaptor protein; SLAP;
XX myeloid malignancy; acute myelogenous leukemia; autoimmune disorder;
XX immunosuppression; myeloproliferative disorder; breast cancer.

XX Mus sp.

XX MO200242452-A2.

XX 30-MAY-2002.

XX 26-NOV-2001; 2001MO-CA01662.

XX 27-NOV-2000; 2000CA-2324663.

XX (HOSP-) HOSPITAL FOR SICK CHILDREN.

XX Meglade JC, Loreto MP;

XX WPI; 2002-566564/60.

XX N-PSDB; AAL44087.

XX New isolated modulator of antigen receptor signalling protein or its
XX fragment, useful for treating malignant disorders such as myeloid
XX malignancies, autoimmune disorders and myeloproliferative disorders -

XX Claim 7; Fig 1A; 110pp; English.

XX The invention comprises the amino acid and coding sequences of modulator
XX of antigen receptor signalling (MARS) proteins. The MARS protein is a
XX putative tumour suppressor gene and exhibits structural and sequence

CC similarity to the Scr-like adaptor protein (SAP). The MARS DNA and
 CC protein sequences of the invention are useful for the treatment of
 CC myeloid malignancies (e.g. acute myelogenous leukaemia) autoimmune
 CC disorders, immunosuppression, myeloproliferative disorders and
 CC malignancies related to the de-regulation of tyrosine kinases (e.g.
 CC breast cancer). The present amino acid sequence represents a mouse MARS
 CC protein.

XX
 SQ Sequence 259 AA;

Query Match 84.0%; Score 242; DB 23; Length 259;
 Best Local Similarity 85.5%; Pred. No. 2, 2e-23;
 Matches 47; Conservative 4; Mismatches 4; Indels 0; Gaps 0;

QY 2 TAVAGSFPAGCPAELSLRLGEPITTSYSDGDMWTVLSVSGREYNIPSYHAKV 56
 DB 35 TAVAGSFPAGCPAELSLRLGEPITTSYSDGDMWTVLSVSGREYNIPSYHAKV 89

RESULT 6
 ABG05994

XX ABG05994 standard; Protein, 70 AA.

AC ABG05994;

DT 13-FEB-2002 (First entry)

XX Novel human diagnostic protein #5985.

XX Human; chromosome mapping; gene mapping; gene therapy; forensic;
 KW food supplement; medical imaging; diagnostic; genetic disorder.

OS Homo sapiens.

PN W0200175067-A2.

XX 11-OCT-2001.

PF 30-MAR-2001; 2001WO-US08631.

XX 31-MAR-2000; 2000US-0540217.

PR 23-AUG-2000; 2000US-0649167.

XX (HYSE-) HYSEQ INC.

PI Dmanac RT, Liu C, Tang YT,

XX WPI; 2001-639362/73.

DR N-PSDB; AAS70181.

XX New isolated polynucleotide and encoded polypeptides, useful in
 PT diagnosis, forensic, gene mapping, identification of mutations
 PT responsible for genetic disorders or other traits and to assess
 PT biodiversity -

XX Claim 20; SEQ ID No 36353; 103pp; English.

CC The invention relates to isolated polynucleotide (I) and
 CC polypeptide (II) sequences. (I) is useful as hybridisation probes,
 CC polymerase chain reaction (PCR) primers, oligomers, and for chromosome
 CC and gene mapping, and in recombinant production of (II). The
 CC polynucleotides are also used in diagnostics as expressed sequence tags
 CC for identifying expressed genes. (II) is useful in gene therapy techniques
 CC to restore normal activity of (II) or to treat disease states involving
 CC (II). (II) is useful for generating antibodies against it, detecting or
 CC quantitating a polypeptide in tissue, as molecular weight markers and as
 CC a food supplement. (II) and its binding partners are useful in medical
 CC disorders involving aberrant protein expression or biological activity.
 CC The polypeptide and polynucleotide sequences have applications in
 CC diagnostics, forensic, gene mapping, identification of mutations
 CC responsible for genetic disorders or other traits to assess biodiversity
 CC and to produce other types of data and products dependent on DNA and

CC amino acid sequences. ABG00010-ABG30377 represent novel human
 CC diagnostic amino acid sequences of the invention.
 CC Note: The sequence data for this patent did not appear in the printed
 CC specification, but was obtained in electronic format directly from WIPO
 CC at ftp.wipo.int/pub/published_pct_sequences.

XX
 SQ Sequence 70 AA;

Query Match 51.4%; Score 148; DB 22; Length 70;
 Best Local Similarity 96.4%; Pred. No. 8, 4e-12;
 Matches 27; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 29 SEDGDMWTVLSVSGREYNIPSYHAKV 56
 DB 6 SKDGDWMTVLSVSGREYNIPSYHAKV 33

RESULT 7
 AAU31598

XX AAU31598 standard; Protein, 395 AA.

AC AAU31598;

DT 18-DEC-2001 (First entry)

XX Novel human secreted protein #2089.

XX Human; vaccination; gene therapy; nutritional supplement;
 KW stem cell proliferation; haematopoiesis; nerve tissue regeneration;
 KW immune suppression; immune stimulation; anti-inflammatory; leukaemia.

OS Homo sapiens.

PN W0200179449-A2.

XX 25-OCT-2001.

PF 16-APR-2001; 2001WO-US08656.

XX 18-APR-2000; 2000US-0552929.

PR 26-JAN-2001; 2001US-0770160.

XX (HYSE-) HYSEQ INC.

PI Tang YT, Liu C, Dmanac RT;

XX WPI; 2001-611725/70.

XX Nucleic acids encoding a range of human polypeptides, useful in genetic
 PT vaccination, testing and therapy -
 PT Claim 20; Page 464-465; 765pp; English.

CC The invention relates to novel human secreted polypeptides. The
 CC polypeptides and antibodies to the polypeptides are useful for
 CC determining the presence of or predisposition to a disease associated
 CC with altered levels of polypeptide. The polypeptides are also useful for
 CC identifying agents (agonists and antagonists) that bind to them. Cells
 CC expressing the proteins are useful for identifying a therapeutic agent
 CC for use in treatment of a pathology related to aberrant expression or
 CC physiological interactions of the polypeptide. Vectors comprising
 CC the nucleic acids encoding the polypeptides and cells genetically
 CC engineered to express them are also useful for producing the proteins.
 CC The proteins are useful in genetic vaccination, testing and
 CC therapy, and can be used as nutritional supplements. They may be used to
 CC increase stem cell proliferation; to regulate haematopoiesis; and in
 CC bone, cartilage, tendon and/or nerve tissue growth or regeneration;
 CC immune suppression and/or stimulation; as anti-inflammatory agents; and
 CC in treatment of leukaemia. AAU29510-AAU3304 represent the amino acid
 CC sequences of novel human secreted proteins of the invention.

XX
 SQ Sequence 395 AA;

Query Match 50.3%; Score 145; DB 22; Length 395;
Best Local Similarity 100.0%; Pred. No. 1.9e-10;
Matches 30; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 ATNALGSPAGPABSLRLGEPITVSE 30
DB 366 ATNALGSPAGPABSLRLGEPITVSE 395

RESULT 8

AAU31072
ID AAU31072 standard; Protein; 315 AA.

AC AAU31072;

DT 18-DEC-2001 (first entry)

DE Novel human secreted protein #1563.

KW Human; vaccination; gene therapy; nutritional supplement;
KW stem cell proliferation; haematopoiesis; nerve tissue regeneration;
KW immune suppression; immune stimulation; anti-inflammatory; leukaemia.

OS Homo sapiens.

PN WO200179449-A2.

PD 25-OCT-2001.

PF 16-APR-2001; 2001MO-US08656.

PR 18-APR-2000; 2000US-0552929.

PR 26-JAN-2001; 2001US-0770160.

PA (HYSE-) HYSEQ INC.

PI Tang YF, Liu C, Drmanac RT,

DR WPI; 2001-611725/70.

PT Nucleic acids encoding a range of human polypeptides, useful in genetic

PT vaccination, testing and therapy -

XX Claim 20; Page 399; 765pp; English.

XX The invention relates to novel human secreted polypeptides. The
CC polypeptides and antibodies to the polypeptides are useful for
CC determining the presence of or predisposition to a disease associated
CC with altered levels of polypeptide. The polypeptides are also useful for
CC identifying agents (agonists and antagonists) that bind to them. Cells
CC expressing the proteins are useful for identifying a therapeutic agent
CC for use in treatment of a pathology related to aberrant expression or
CC physiological interactions of the polypeptide. Vectors genetically
CC engineered to express them are also useful for producing the proteins.
CC The proteins are useful in genetic vaccination, testing and
CC therapy, and can be used as nutritional supplements. They may be used to
CC increase stem cell proliferation; to regulate haematopoiesis, and in
CC bone, cartilage, tendon and/or nerve tissue growth or regeneration;
CC immune suppression and/or stimulation; as anti-inflammatory agents; and
CC in treatment of leukaemias. AAU29510-AAU3304 represent the amino acid
CC sequences of novel human secreted proteins of the invention.

SO Sequence 315 AA;

QY Query Match 34.4%; Score 99; DB 22; Length 315;
Best Local Similarity 37.3%; Pred. No. 0.00015;
Matches 19; Conservative 11; Mismatches 21; Indels 0; Gaps 0;

QY 6 LGSFPAGPABSLRLGEPITVSEDDGDMWTVLSVSGREYNIPSYHAKV 56
DB 68 LGSFPAGPABSLRLGEPITVSEDDGDMWTVLSVSGREYNIPSYHAKV 118

RESULT 9
AAV49420
ID AAV49420 standard; Protein; 509 AA.

AC AAV49420;

DT 13-MAR-2000 (first entry)

DE PKA substrate, Src-family protein.

KW Protein kinase A; PKA; PKA signaling pathway; phosphorylation; cancer;
KW kinase substrate; immunosuppressive disorder; proliferative disease;
KW HIV infection; AIDS; immunodeficiency; autoimmune disease;
KW systemic lupus erythematosus; Src-family.

OS Homo sapiens.

PN WO9962315-A2.

PD 02-DEC-1999.

PF 27-MAY-1999; 99WO-GB01680.

PR 27-MAY-1998; 98NO-0002419.

PR 30-DEC-1998; 98US-0114240.

PA (LAUR-) LAURAS AS.

PI (JONES) JONES E L.

PI Hansson V, Levy FO, Mustelin T, Skalhogg BS, Sundvold V, Tasken K;

DR Vang T, Altman A, Munshi A;

DR WPI; 2000-086801/07.

DR N-PSDB; AA246491.

PT Altering the activity of protein kinase signaling pathways, used for

PT treating immunosuppressive disorders, e.g. AIDS, proliferative

PT disorders, e.g. cancers or autoimmune diseases -

XX Claim 23; Page 95-96; 111pp; English.

XX The invention provides a novel method of altering the activity of the
CC protein kinase A (PKA) signaling pathway in a cell that comprises
CC altering the extent of phosphorylation of one or more PKA substrates, or
CC kinase substrates downstream in the PKA signaling pathway. Pharmaceutical
CC compositions containing a nucleic acid molecule that encodes a PKA
CC substrate, or fragment, precursor or functionally equivalent variant,
CC where the sequence is modified to alter its susceptibility to
CC phosphorylation by PKA can be used for treating a disorder exhibiting
CC abnormal PKA signaling activity, immunosuppressive disorders or
CC proliferative diseases. They can be used for treating e.g. HIV
CC infection, AIDS, common variable immunodeficiency or cancers. Conditions
CC in which upregulation of the PKA pathway is required, such as autoimmune
CC disease, e.g. systemic lupus erythematosus, may also be treated. The
CC present sequence represents a PKA substrate, wherein the substrate is in
CC the Src-family, preferably Lck, Fyn, Src, Yes, Fgr, Lyn, Hck Blk, Yrk,
CC c-trl, Fyk, Src-1 or Src-2.

SO Sequence 509 AA;

QY Query Match 33.3%; Score 96; DB 21; Length 509;
Best Local Similarity 38.5%; Pred. No. 0.00066;
Matches 20; Conservative 8; Mismatches 24; Indels 0; Gaps 0;

QY 4 VALGSPAGPABSLRLGEPITVSEDDGDMWTVLSVSGREYNIPSYHAKV 55
DB 67 VALGSPAGPABSLRLGEPITVSEDDGDMWTVLSVSGREYNIPSYHAKV 118

RESULT 10
ABB57957
ID ABB57957 standard; Protein; 517 AA.

XX AC ABB57957;
 XX XX
 DT 26-MAR-2002 (first entry)
 XX XX
 DE Drosophila melanogaster polypeptide SEQ ID NO 663.
 XX XX
 KM Drosophila; developmental biology; cell signalling; insecticide;
 KM pharmaceutical.
 XX OS Drosophila melanogaster.
 XX PN MO200171042-A2.
 XX XX
 PD 27-SEP-2001.
 XX XX
 PP 23-MAR-2001; 2001MO-US09231.
 XX XX
 PR 23-MAR-2000; 2000US-191637P.
 XX PR 11-JUL-2000; 2000US-0614150.
 XX XX
 PA (PEKE) PE CORP NY.
 XX XX
 PI Venter JC, Adams M, Li PWD, Myers EW;
 XX XX
 DR WPI; 2001-656860/75.
 DR N-PSDB; ABL02060.
 XX XX
 PT New isolated nucleic acid detection reagent for detecting 1000 or more
 PT genes from Drosophila and for elucidating cell signalling and cell-cell
 PT interactions -
 XX XX
 PS Disclosure; SEQ ID NO 663; 21pp + Sequence Listing; English.
 CC The invention relates to an isolated nucleic acid detection reagent
 CC capable of detecting 1000 or more genes from Drosophila. The invention is
 CC useful in developmental biology and in elucidating cell signalling and
 CC cell-cell interactions in higher eukaryotes for the development of
 CC insecticides, therapeutic and pharmaceutical drugs. The invention
 CC discloses genomic DNA sequences (AB16175-AB130511), expressed DNA
 CC sequences (AB16175-AB16175) and the encoded proteins
 CC (AB16175-AB16175).
 CC The sequence data for this patent did not form part of the printed
 CC specification, but was obtained in electronic format directly from WIPO
 CC at ftp.wipo.int/pub/published_pct_sequences.
 XX XX
 SQ Sequence 517 AA;
 Query Match 31.8%; Score 91.5; DB 22; Length 517;
 Best Local Similarity 42.6%; Pred. No. 0.0027;
 Matches 23; Conservative 9; Mismatches 21; Indels 1; Gaps 1;
 QY 4 VALGFPAGPAEISLRIGEPITVSE-DGDMWTVLSEVSGREYNIPSVYAKV 56
 Db 69 VALYDYDARTDEDLFRKGEHLINDTGDWMVLARSKTRSGYIPSNVAKL 122
 RESULT 11
 ABB84051
 ID ABB84051 standard; protein; 61 AA.
 XX AC ABB84051;
 XX XX
 DT 04-SEP-2002 (first entry)
 XX XX
 DE Human protein fragment capable of inactivating HIV Nef protein.
 XX XX
 KM Nef protein; fusion protein; virucide; anti-HIV; accessory protein;
 KM pathogenicity; diagnosis; AIDS; human.
 XX OS Homo sapiens.
 XX PN DE10109532-C1.

XX PD 13-JUN-2002.
 XX XX
 PP 28-FEB-2001; 2001DE-1009532.
 XX XX
 PR 28-FEB-2001; 2001DE-1009532.
 XX XX
 PA (GEVE/) GEYER M.
 PA (PACK/) FACKLER O.
 XX XX
 PI Geyer M;
 XX XX
 DR WPI; 2002-418264/45.
 XX XX
 PT New fusion protein that blocks Nef protein, useful for treatment or
 PT diagnosis of acquired immune deficiency syndrome, has high specificity
 PT and affinity -
 XX XX
 PS Claim 3; Page 8-9; 22pp; German.
 CC This invention describes a novel fusion protein for blocking the Nef
 CC protein of human immune deficiency virus (HIV) which comprises: (i)
 CC protein domain 1 that binds to a di-leucine (LL) motif; (ii) a
 CC linker between protein domains 1 and 2. The products of the invention
 CC have virucide and anti-HIV activity and are capable of neutralising Nef,
 CC an accessory protein essential for pathogenicity of HIV-1. The fusion
 CC protein of the invention comprises the LL domain of the beta-subunit of
 CC the adapter-protein complex Ap-1 and the PxxP binding SH3 domain of
 CC tyrosine kinase Hck, linked through a 60 amino acid peptide. The products
 CC of the invention are used for in vitro diagnosis of AIDS and for
 CC treatment of AIDS. The LL and PxxP motifs are specific for Nef, which,
 CC unlike HIV protease, has no human homologue, so the fusion protein (which
 CC binds Nef with very high affinity) should cause essentially no side
 CC effects. This sequence represents a human derived protein fragment used
 CC in the construction of the fusion protein of the invention and which
 CC contains a PxxP-motif binding domain useful to the invention.
 XX XX
 SQ Sequence 61 AA;
 Query Match 31.2%; Score 90; DB 23; Length 61;
 Best Local Similarity 35.8%; Pred. No. 0.0028;
 Matches 19; Conservative 12; Mismatches 22; Indels 0; Gaps 0;
 QY 4 VALGFPAGPAEISLRIGEPITVSE-DGDMWTVLSEVSGREYNIPSVYAKV 56
 Db 7 VALYDYDARTDEDLFRKGEHLINDTGDWMVLARSKTRSGYIPSNVAKV 59
 RESULT 12
 ABB84052
 ID ABB84052 standard; protein; 471 AA.
 XX AC ABB84052;
 XX XX
 DT 04-SEP-2002 (first entry)
 XX XX
 DE Rac/human fusion protein capable of inactivating HIV Nef protein.
 XX XX
 KM Nef protein; fusion protein; virucide; anti-HIV; accessory protein;
 KM pathogenicity; diagnosis; AIDS; rac; human.
 XX OS Ratius sp.
 XX OS Homo sapiens.
 XX OS Synthetic.
 XX XX
 PH Key
 PH Region
 FT 1..349
 FT /note= "Rat derived region"
 FT 427..471
 FT /note= "Human derived region"
 XX XX
 PN DE10109532-C1.

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XX 13-JUN-2002.
PD
XX
XX 28-FEB-2001; 2001DE-1009532.
PF
XX 28-FEB-2001; 2001DE-1009532.
PR
XX (GEVE/) GEYER M.
PA (PACK/) FACKLER O.
XX
PI Geyer M;
DR WPI; 2002-418264/45.
XX
XX New fusion protein that blocks Nef protein, useful for treatment or
PT diagnosis of acquired immune deficiency syndrome, has high specificity
XX and affinity
XX
PS Claim 4; Page 9-11; 22pp; German.
XX
XX This invention describes a novel fusion protein for blocking the Nef
CC protein of human immune deficiency virus (HIV) which comprises: (i) a
CC protein domain 1 that binds to a di-leucine (LL) motif; (ii) a
CC protein domain 2 that binds to a PxxP motif; and (iii) a polypeptide
CC linker between protein domains 1 and 2. The products of the invention
CC have virucide and anti-HIV activity and are capable of neutralising Nef,
CC an accessory protein essential for pathogenicity of HIV-1. The fusion
CC protein of the invention comprises the LL domain of the beta-subunit of
CC the adapter-protein complex Ap-1 and the PxxP binding SH3 domain of
CC tyrosine kinase Hck, linked through a 60 amino acid peptide. The products
CC of the invention are used for in vitro diagnosis of AIDS and for
CC treatment of AIDS. The LL and PxxP motifs are specific for Nef, which,
CC unlike HIV protease, has no human homologue, so the fusion protein (which
CC binds Nef with very high affinity) should cause essentially no side
CC effects. This sequence represents a fusion protein composed of a rat
CC protein fragment which contains a dileucine (LL) motif and a human
CC protein fragment containing a PxxP-motif binding domain useful to the
CC invention.
XX
SQ Sequence 471 AA;
Query Match 31.2%; Score 90; DB 23; Length 471;
Best Local Similarity 35.8%; Pred. No. 0.0038;
Matches 19; Conservative 12; Mismatches 22; Indels 0; Gaps 0;
QY 4 VALGSPAGGPAELSLRGEPLTIVSEDDGMWTVLSVSGREYNISYHAKV 56
DB 417 VALYDYEAHHEHDLSPFGKQDMVTVLESGEWMKARSLATRKRGYIPSNVAVY 469
RESULT 13
AAB99332
ID AAB99332 standard; Protein; 505 AA.
XX
AC AAB99332;
XX
XX 23-AUG-2001 (first entry)
DT
XX
XX Human tyrosine kinase Hck protein sequence SEQ ID NO:11.
DE
XX
XX Human; tyrosine kinase Hck binding protein; tyrosine kinase; Hck;
KW tumour lethal factor; tumour necrosis factor alpha; apoptosis; HSB-1;
KW Hck signal transduction; human immunodeficiency virus; HIV infection;
XX anticancer.
XX
XX Homo sapiens.
OS
XX
XX WO200132869-A1.
PN
XX
XX 10-MAY-2001.
PD
XX
XX 26-OCT-2000; 2000WO-JP07500.
PF
XX

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PR 29-OCT-1999; 99JP-0309957.
XX
XX (SSSE ) SSP CO LTD.
PA
XX Taniyama T, Narita T;
PI
XX WPI; 2001-316440/33.
DR
XX
XX New proteins which bind to human tyrosine kinase Hck for promotion of
PT apoptosis and for the elucidation of the mechanism of Hck signal
PT transduction
XX
XX Example 1; Page 33-35; 45pp; Japanese.
PS
XX
XX The present invention describes a protein, designated HSB-1, which binds
CC to human tyrosine kinase Hck. Also described are: (1) nucleic acids
CC encoding the protein and its derivatives; (2) recombinant vectors
CC containing the nucleic acids; and (3) host cells transformed by the
CC vectors and expressing the protein. HSB-1 has cytoprotective activity, binds
CC tyrosine kinase, enhances tumour necrosis factor alpha and promotes
CC apoptosis. HSB-1 proteins are used for the elucidation of the mechanism
CC of Hck signal transduction and of the role of Hck in human
CC immunodeficiency virus (HIV) infection. They can be used for the
CC treatment of infections and other diseases with which Hck is associated.
CC They promote the anticancer activity of tumour necrosis factor alpha.
CC The present sequence represents the human tyrosine kinase Hck protein,
CC which is used in an example from the present invention.
XX
SQ Sequence 505 AA;
Query Match 31.2%; Score 90; DB 22; Length 505;
Best Local Similarity 35.8%; Pred. No. 0.0041;
Matches 19; Conservative 12; Mismatches 22; Indels 0; Gaps 0;
QY 4 VALGSPAGGPAELSLRGEPLTIVSEDDGMWTVLSVSGREYNISYHAKV 56
DB 63 VALYDYEAHHEHDLSPFGKQDMVTVLESGEWMKARSLATRKRGYIPSNVAVY 115
RESULT 14
AAB84055
ID AAB84055 standard; protein; 513 AA.
XX
AC AAB84055;
XX
XX 04-SEP-2002 (first entry)
DT
XX
XX Rat/human fusion protein capable of inactivating HIV Nef protein.
DE
XX
XX Nef protein; fusion protein; virucide; anti-HIV; accessory protein;
KW pathogenicity; diagnosis; AIDS; rat; human.
XX
XX Rattus sp.
OS
XX Homo sapiens.
OS
XX Synthetic.
XX
XX Key 1.349 Location/Qualifiers
FH Region /note="Rat derived region"
FT 407..426
FT Region /note="human ubiquitination signal"
FT 446..490
FT Region /note="Human derived region"
FT 491..513
FT Region /note="Human farneylelation signal"
XX
XX DE10109532-C1.
PN
XX
XX 13-JUN-2002.
PD
XX
XX 28-FEB-2001; 2001DE-1009532.
PF
XX
XX 28-FEB-2001; 2001DE-1009532.
PR

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XX (GEYE/) GEYER M.
 PA (PACK/) FACKLER O.
 XX
 PI Geyer M.
 XX WPI, 2002-418264/45.
 DR
 XX New fusion protein that blocks Nef protein, useful for treatment or
 PT diagnosis of acquired immune deficiency syndrome, has high specificity
 PT and affinity -
 XX
 PS Claim 9, Page 11-13; 22pp; German.

CC This invention describes a novel fusion protein for blocking the Nef
 CC protein of human immune deficiency virus (HIV) which comprises: (i)
 CC protein domain 1 that binds to a di-leucine (LL) motif; (ii) a
 CC protein domain 2 that binds to a PXXP motif; and (iii) a polypeptide
 CC linker between protein domains 1 and 2. The products of the invention
 CC have virucide and anti-HIV activity and are capable of neutralizing Nef,
 CC an accessory protein essential for pathogenicity of HIV-1. The fusion
 CC protein of the invention comprises the LL domain of the beta-subunit of
 CC the adapter-protein complex AP-1 and the PXXP binding SH3 domain of
 CC tyrosine kinase Hck, linked through a 60 amino acid peptide. The products
 CC of the invention are used for in vitro diagnosis of AIDS and for
 CC treatment of AIDS. The LL and PXXP motifs are specific for Nef, which,
 CC unlike HIV protease, has no human homologue, so the fusion protein (which
 CC binds Nef with very high affinity) should cause essentially no side
 CC effects. This sequence represents a fusion protein composed of a rat
 CC protein fragment which contains a dileucine (LL) motif and a human
 CC protein fragment containing a PXXP-motif binding domain, a farnesylation
 CC signal and a ubiquitination signal useful to the invention.

CC Sequence 513 AA;

Query Match 31.2%; Score 90; DB 23; Length 513;
 Best Local Similarity 35.8%; Pred. No. 0.0042;
 Matches 19; Conservative 12; Mismatches 22; Indels 0; Gaps 0;

Oy 4 VALGSPAGPAELSLRLGEPITV-SEGDGMWTVLSEVSGREYNIPSVHAKV 56
 Db 436 VALDYEAHHEHDLSPKQDQWVLESGEWMKARSLATRKGYIPSNYVARV 488

RESULT 15

ID ABB58710 standard; Protein; 2415 AA.

XX ABB58710;

XX 26-MAR-2002 (first entry)

XX Drosophila melanogaster polypeptide SEQ ID NO 2922.

XX Drosophila; developmental biology; cell signalling; insecticide;

XX pharmaceutical.

XX Drosophila melanogaster.

XX WO200171042-A2.

XX 27-SEP-2001.

XX 23-MAR-2001; 2001WO-US09231.

XX 23-MAR-2000; 2000US-191637P.

XX 11-JUL-2000; 2000US-0614150.

XX (PEKE & PE CORP NY.

XX Venter JC, Adams M, Li PMD, Myers EW;
 DR WPI, 2001-656860/75.

DR N-PSDB; ABL02813.

XX New isolated nucleic acid detection reagent for detecting 1000 or more
 PT genes from Drosophila and for elucidating cell signalling and cell-cell
 PT interactions -

XX Disclosure; SEQ ID NO 2922; 21pp + Sequence Listing; English.

CC The invention relates to an isolated nucleic acid detection reagent
 CC capable of detecting 1000 or more genes from Drosophila. The invention is
 CC useful in developmental biology and in elucidating cell signalling and
 CC cell-cell interactions in higher eukaryotes for the development of
 CC insecticides, therapeutic and pharmaceutical drugs. The invention
 CC discloses genomic DNA sequences (ABL16176-ABL10511), expressed DNA
 CC sequences (ABL01840-ABL16175) and the encoded proteins
 CC (ABB57737-ABB72072).
 CC The sequence data for this patent did not form part of the printed
 CC specification, but was obtained in electronic format directly from WIPO
 CC at ftp.wipo.int/pub/published_pct_sequences.

XX Sequence 2415 AA;

Query Match 31.1%; Score 89.5; DB 22; Length 2415;
 Best Local Similarity 35.2%; Pred. No. 0.036;
 Matches 19; Conservative 15; Mismatches 17; Indels 3; Gaps 2;

Oy 4 VALGSPAGPAELSLRLGEPITV-SEGDGMWTVLSEVSGREYNIPSVHAKV 56
 Db 976 VALDYERKSPREVSMMKQDVITLNSNNKDMWKKV--EVNDRQGFVPAAYIKKI 1027

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